



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/009,792A

DATE: 04/03/2003

TIME: 14:26:35

Input Set : A:\HYLEE60.001.TXT

Output Set: N:\CRF4\04032003\J009792A.raw

C--> 4 <110> APPLICANT: LEE, Sang-Yup
5 JEONG, Ki-Jun
7 <120> TITLE OF INVENTION: ESCHERICHIA COLI STRAIN SECRETING HUMAN
8 GRANULOCYTE COLONY STIMULATING FACTOR (G-CSF)
11 <130> FILE REFERENCE: HYLEE60.001APC
13 <140> CURRENT APPLICATION NUMBER: US 10/009,792A
14 <141> CURRENT FILING DATE: 2002-10-29
16 <150> PRIOR APPLICATION NUMBER: PCT/KR01/00549
17 <151> PRIOR FILING DATE: 2001-03-31
19 <150> PRIOR APPLICATION NUMBER: KR 10-2000-0017052
20 <151> PRIOR FILING DATE: 2000-03-31
22 <160> NUMBER OF SEQ ID NOS: 27
24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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27 <211> LENGTH: 13
28 <212> TYPE: PRT
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Oligopeptide
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37 1 5 10
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41 <211> LENGTH: 29
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43 <213> ORGANISM: Artificial Sequence
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51 <210> SEQ ID NO: 3
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53 <212> TYPE: DNA
54 <213> ORGANISM: Artificial Sequence
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63 <211> LENGTH: 50
64 <212> TYPE: DNA
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ENTERED

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68 <223> OTHER INFORMATION: Primer
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76 <213> ORGANISM: Artificial Sequence
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84 <210> SEQ ID NO: 6
85 <211> LENGTH: 50
86 <212> TYPE: DNA
87 <213> ORGANISM: Artificial Sequence
89 <220> FEATURE:
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96 <211> LENGTH: 48
97 <212> TYPE: DNA
98 <213> ORGANISM: Artificial Sequence
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103 <400> SEQUENCE: 7
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106 <210> SEQ ID NO: 8
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117 <210> SEQ ID NO: 9
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129 <211> LENGTH: 33
130 <212> TYPE: DNA
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137 qqaattcaca tdtttaagtt taaaaaagaaa ttc 33
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152 <212> TYPE: DNA
153 <213> ORGANISM: Artificial Sequence
155 <220> FEATURE:
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158 <400> SEQUENCE: 12
159 gcaaccgcct ctgcaactcc gttagggtcca gcc 33
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162 <211> LENGTH: 33
163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: Primer
169 <400> SEQUENCE: 13
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172 <210> SEQ ID NO: 14
173 <211> LENGTH: 36
174 <212> TYPE: DNA
175 <213> ORGANISM: Artificial Sequence
177 <220> FEATURE:
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180 <400> SEQUENCE: 14
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184 <211> LENGTH: 39
185 <212> TYPE: DNA
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191 <400> SEQUENCE: 15
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195 <211> LENGTH: 41
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206 <211> LENGTH: 507
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208 <213> ORGANISM: Homo sapiens
210 <400> SEQUENCE: 17
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212 agtgcactct ggacagtgc ggaagccacc cccctgggcc ctgccagctc cctgccccag 120
213 agcttctctgc tcaagtgtt agagcaagtg aggaagatcc agggcgatgg cgcagcgctc 180
214 caggagaagc tggcaggctg cttgagccaa ctccatagcg gcttttctt ctaccagggg 240
215 ctctgcagg ccttggagg gatctcccc gagttgggtc ccaccttga cacactgcag 300
216 ctggacgtcg ccgaatttc caccaccatc tggcagcaga tggagaact gggaatggcc 360
217 cctgccccgc agccaccca gggtgccatg ccggccttcg cctctgttt ccagcgccgg 420
218 gcaggagggg tctagtgtc ctcccatctg cagagcttc tggaggtgtc gtaccgcgtt 480
219 ctacgccacc ttgccagcc ctaataa 507
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222 <211> LENGTH: 615
223 <212> TYPE: DNA
224 <213> ORGANISM: Homo sapiens
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229 agcttctctgc tcaagtgtt agagcaagtg aggaagatcc agggcgatgg cgcagcgctc 180
230 caggagaagc tgtgtgccac ctacaagctg tgccacccc aggagctggt getgctcgga 240
231 cactctctgg gcateccctg ggctccctg agcagctgcc ccagccagge cctgcagctg 300
232 gcaggctgct tgagccaact ccatagcggc ctttctctt accaggggct cctgcaggcc 360
233 ctggaaggga tctccccga gttgggtccc accttgaca cactgcagct ggacgtcgcc 420
234 gactttgcca ccaccatctg gcagcagatg gaaqaactgg gaatggccc tgcctgcag 480
235 cccacccagg gtgccatgcc ggccttcgcc tctgttttc agcgccgggc aggaggggtc 540
236 ctagtgtcct cccatctgca gagcttctct gaggtgtcgt accgcgttct accccacctt 600
237 gccagccct aataa 615
239 <210> SEQ ID NO: 19
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241 <212> TYPE: PRT
242 <213> ORGANISM: Homo sapiens
244 <400> SEQUENCE: 19
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246 1 5 10 15
247 Cys Leu Glu Gln Val Arg Lys Ile Gln Gly Asp Gly Ala Ala Leu Gln
248 20 25 30
249 Glu Lys Leu Cys Ala Thr Tyr Lys Leu Cys His Pro Glu Glu Leu Val
250 35 40 45
251 Leu Leu Gly His Ser Leu Gly Ile Pro Trp Ala Pro Leu Ser Ser Cys
252 50 55 60
253 Pro Ser Gln Ala Leu Gln Leu Ala Gly Cys Leu Ser Gln Leu His Ser
254 65 70 75 80
255 Gly Leu Phe Leu Tyr Gln Gly Leu Leu Gln Ala Leu Glu Gly Ile Ser
256 85 90 95
257 Pro Glu Leu Gly Pro Thr Leu Asp Thr Leu Gln Leu Asp Val Ala Asp
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258          100          105          110
259 Phe Ala Thr Thr Ile Trp Gln Gln Met Glu Glu Leu Gly Met Ala Pro
260          115          120          125
261 Ala Leu Gln Pro Thr Gln Gly Ala Met Pro Ala Phe Ala Ser Ala Phe
262          130          135          140
263 Gln Arg Arg Ala Gly Gly Val Leu Val Ala Ser His Leu Gln Ser Phe
264 145          150          155          160
265 Leu Glu Val Ser Tyr Arg Val Leu Arg His Leu Ala Gln Pro
266          165          170
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270 <211> LENGTH: 531
271 <212> TYPE: DNA
272 <213> ORGANISM: Homo sapiens
274 <400> SEQUENCE: 20
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276 caagtgagga agatccaggg cgtggcgca gcgtccagg agaagctgtg tgccacctac 120
277 aagctgtgcc accccgagga gctggtgctg ctgggacaact ctctgggcat cccctgggct 180
278 cccctgagca gctgccccag ccaggccctg cagctggcag gtgctttgag ccaactccat 240
279 agcggccttt tctctacca ggggctcctg caggccctgg aagggatctc ccccgagttg 300
280 ggtccacct tggacacaact gcagctggac gtgcgcgact ttgccaccac catctggcag 360
281 cagatggaag aactgggaat ggcctctgcc ctgcagccca ccagggtgc catgccggcc 420
282 ttgcctctg cttccagcg ccgggcagga ggggtcctag ttgcctccca tetgcagagc 480
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285 <210> SEQ ID NO: 21
286 <211> LENGTH: 175
287 <212> TYPE: PRT
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290 <400> SEQUENCE: 21
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293 Lys Cys Leu Glu Gln Val Arg Lys Ile Gln Gly Asp Gly Ala Ala Leu
294 20 25 30
295 Gln Glu Lys Leu Cys Ala Thr Tyr Lys Leu Cys His Pro Glu Glu Leu
296 35 40 45
297 Val Leu Leu Gly His Ser Leu Gly Ile Pro Trp Ala Pro Leu Ser Ser
298 50 55 60
299 Cys Pro Ser Gln Ala Leu Gln Leu Ala Gly Cys Leu Ser Gln Leu His
300 65 70 75 80
301 Ser Gly Leu Phe Leu Tyr Gln Gly Leu Leu Gln Ala Leu Glu Gly Ile
302 85 90 95
303 Ser Pro Glu Leu Gly Pro Thr Leu Asp Thr Leu Gln Leu Asp Val Ala
304 100 105 110
305 Asp Phe Ala Thr Thr Ile Trp Gln Gln Met Glu Glu Leu Gly Met Ala
306 115 120 125
307 Pro Ala Leu Gln Pro Thr Gln Gly Ala Met Pro Ala Phe Ala Ser Ala
308 130 135 140
309 Phe Gln Arg Arg Ala Gly Gly Val Leu Val Ala Ser His Leu Gln Ser
310 145 150 155 160
311 Phe Leu Glu Val Ser Tyr Arg Val Leu Arg His Leu Ala Gln Pro
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VERIFICATION SUMMARY

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date